



## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>6</sup> :</b> <b>C12P 19/34, C12Q 1/68, 1/70, G01N 33/53, C07H 21/04, C07K 16/00</b>	<b>A1</b>	<b>(11) International Publication Number:</b> <b>WO 97/03207</b> <b>(43) International Publication Date:</b> 30 January 1997 (30.01.97)
<b>(21) International Application Number:</b> PCT/US96/11619 <b>(22) International Filing Date:</b> 12 July 1996 (12.07.96)  <b>(30) Priority Data:</b> 60/002,245      13 July 1995 (13.07.95)      US  <b>(71) Applicant:</b> IMMUNOLOGICAL ASSOCIATES OF DENVER [US/US]; Suite 330, 101 University Boulevard, P.O. Box 61159, Denver, CO 80206 (US).  <b>(72) Inventor:</b> GERDES, John, C.; 375 Steele Street, Denver, CO 80206 (US).  <b>(74) Agent:</b> BERNARD, Julie, L.; 9000 E. Inspiration Drive, Parker, CO 80134 (US).		<b>(81) Designated States:</b> AU, CA, JP, European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).  <b>Published</b> <i>With international search report.</i>
<b>(54) Title:</b> DETECTION OF AMPLIFIED NUCLEIC ACID SEQUENCES USING BIFUNCTIONAL HAPTENIZATION AND DYED MICROPARTICLES  <b>(57) Abstract</b>  The invention describes an assay for detecting amplified target nucleic acid sequences with a visual signal. The sensitivity and specificity of the methodology are based on bifunctional target labeling during the amplification step or subsequent hybridization that generates a bifunctional label. The invention may be used, for example, in the screening of amplicon detection for the purpose of more efficiently screening libraries. The invention is also useful to detect nucleic acid sequences indicative of a genetic defect or contagious disease when used with the appropriate primers, as well as detect the existence of nucleic acid amplification.		

**FOR THE PURPOSES OF INFORMATION ONLY**

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AM	Armenia	GB	United Kingdom	MW	Malawi
AT	Austria	GE	Georgia	MX	Mexico
AU	Australia	GN	Guinea	NE	Niger
BB	Barbados	GR	Greece	NL	Netherlands
BE	Belgium	HU	Hungary	NO	Norway
BF	Burkina Faso	IE	Ireland	NZ	New Zealand
BG	Bulgaria	IT	Italy	PL	Poland
BJ	Benin	JP	Japan	PT	Portugal
BR	Brazil	KE	Kenya	RO	Romania
BY	Belarus	KG	Kyrgyzstan	RU	Russian Federation
CA	Canada	KP	Democratic People's Republic of Korea	SD	Sudan
CF	Central African Republic	KR	Republic of Korea	SE	Sweden
CG	Congo	KZ	Kazakhstan	SG	Singapore
CH	Switzerland	LI	Liechtenstein	SI	Slovenia
CI	Côte d'Ivoire	LK	Sri Lanka	SK	Slovakia
CM	Cameroon	LR	Liberia	SN	Senegal
CN	China	LT	Lithuania	SZ	Swaziland
CS	Czechoslovakia	LU	Luxembourg	TD	Chad
CZ	Czech Republic	LV	Latvia	TG	Togo
DE	Germany	MC	Monaco	TJ	Tajikistan
DK	Denmark	MD	Republic of Moldova	TT	Trinidad and Tobago
EE	Estonia	MG	Madagascar	UA	Ukraine
ES	Spain	ML	Mali	UG	Uganda
FI	Finland	MN	Mongolia	US	United States of America
FR	France	MR	Mauritania	UZ	Uzbekistan
GA	Gabon			VN	Viet Nam

# **DETECTION OF AMPLIFIED NUCLEIC ACID SEQUENCES USING BIFUNCTIONAL HAPTENIZATION AND DYED MICROPARTICLES**

## **FIELD OF INVENTION**

5           This invention relates to the general fields of molecular biology and medical sciences, and specifically to the detection of amplified nucleic acid sequences. This application, thus, describes a method of detecting amplified nucleic acid sequences rapidly and accurately using bifunctional haptenization and dyed microparticles. This application further relates to a  
10       test kit useful for improved detection of amplified nucleic acids.

## **BACKGROUND AND PRIOR ART**

          In current research and development processes involving nucleic acid (DNA and RNA) amplifications, there is a need for rapid, accurate qualitative determination of the results from such experiments.  
15       Traditionally, such determinations require the preparation of gels, for example, agarose, for electrophoresis of amplified nucleic acids and the use of a nucleic acid intercalating agent, for example, ethidium bromide, which is fluorescent under ultraviolet light. Generally, agarose gels are cast by melting a measured amount of agarose in a buffer resulting in a transparent  
20       solution. The melted agarose is poured into a mold and allowed to harden. The mold includes sample wells for adding the nucleic acids to be analyzed. In order to detect the nucleic acid, ethidium bromide is included in the buffer used to prepare the gel or the gel is stained with ethidium bromide after electrophoresis. Samples of the nucleic acids to be analyzed are mixed  
25       with a loading buffer and deposited into the sample wells. The gel is then subjected to a certain applied voltage resulting in the lateral migration of the nucleic acid through the gel. This method requires a power supply and an apparatus for casting and running the gels. Additionally, an ultraviolet light box is required for detecting the nucleic acid, requiring the use of protective  
30       eye wear. Further, the use of ethidium bromide poses an health hazard as it

is a carcinogen, a powerful mutagen and moderately toxic. The qualitative determination process, as described above, is time consuming, requires special precautions for the handling and disposing of ethidium bromide, requires the purchase of specialized equipment, poses a potential health hazard and requires specialized training in the use of the necessary equipment. The sensitivity of the method is low and since equipment is used many times over, the possibility of contamination is also present.

The introduction of nucleic acid probe tests based on hybridization into routine clinical laboratory procedures has been hindered by lack of sensitivity when compared with conventional culture techniques and immunoassays. To increase sensitivity, radioactive labels are used. The problem with the use of such radioactive labels is the laboratory management of radioisotopes and the disposal thereof. To some degree, non-isotopic labels are used to overcome this problem. However, non-isotopic methods do not possess adequate sensitivity and are cumbersome and laborious to run. In order to achieve the sensitivity needed for diagnostics, the incubation times of the current methodologies must be long, taking up to several hours or days. The current non-isotopic methodologies also employ many steps, requiring previous training and expertise.

The ability to amplify nucleic acids from clinical samples has greatly advanced nucleic acid probe technology, providing the sensitivity lacking in earlier versions of non-isotopic assays. The sensitivity afforded by oligonucleotide probe tests utilizing nucleic acid amplification now exceeds that of any other method. However, the method is still time consuming, requiring the use of specialized hybridization techniques and equipment, transfer to solid phases and specialized detection equipment.

The detection of amplified nucleic acids for clinical use largely relies on hybridization of the amplified product with a detection probe that is labeled with a variety of enzymes and luminescent reagents. United States Patent No. 5,374,524 to Miller, describes a nucleic acid probe assay which

combines nucleic acid amplification and solution hybridization using capture and reporter probes. These techniques require multiple reagents, several washing steps, and specialized equipment for detection of the target nucleic acid. Moreover, these techniques are labor intensive and require technicians with expertise in molecular biology.

The use of probes comprised of oligonucleotide sequences bound to microparticles is well known and illustrated in the prior art. The mechanism for attachment of oligonucleotides to microparticles in hybridization assays and for the purification of nucleic acids is also well known. European Patent No. 200 133 describes the attachment of oligonucleotides to water-insoluble particles less than 50 micrometers in diameter used in hybridization assays for the capture of target nucleotides. United States Patent No. 5,387,510 to Wu, describes the use of oligonucleotide sequences covalently bound to microparticles as probes for capturing PCR amplified nucleic acids. United States Patent No. 5,328,825 to Findlay also describes an oligonucleotide linked by way of a protein or carbohydrate to a water-insoluble particle. The oligonucleotide probe is covalently coupled to the microparticle or other solid support. The sensitivity and specificity of the four above-reference patents, each of which is specifically incorporated herein, is based on hybridization of the oligonucleotide probe to the target nucleic acid.

The use of incorporated non-radioactive labels into the amplification reactions for the detection of nucleic acids is also well known in the art. Nucleic acids modified with biotin (United States Patent No. 4,687,732 to Ward *et al.*; European Patent No. 063879; both specifically incorporated herein), digoxin (European Patent No. 173251, specifically incorporated herein) and other haptens have also been used. For example, United States Patent No. 5,344,757 to Graf, specifically incorporated herein, uses a nucleic acid probe containing at least one hapten as a label for hybridization with a complementary target nucleic acid bound to a solid membrane. The

sensitivity and specificity of these assays is based on the incorporation of a single label in the amplification reaction which can be detected using an antibody specific to the label. The usual case involves an antibody conjugated to an enzyme. Furthermore, the addition of substrate generates a colorimetric or fluorescent change which can be detected with an instrument.

Still, the above-described approaches are labor intensive with many steps and washes; require special and costly equipment for the detection of the target nucleic acid; require trained staff; and take a several hours to complete. Several patents have issued which deal with automation of the processes of amplification and subsequent detection of the amplicon. These patents use specialized equipment and are still based on the principle of hybridization and immunoassay technology. For example, European Patent No. 320308, specifically incorporated herein, describes a system for detecting target nucleic acids amplified by the ligase chain reaction.

Automated approaches eliminate the need for specially trained personnel, however, the cost of the equipment is very high and the possibility of contamination still exists since many samples will be processed by the same equipment.

The use of bifunctional labels for detection of an amplified target sequence has been explored using the ligase chain reaction (European Patent No. 320308, *supra*). Upon completion of the amplification by the ligation process, double stranded DNA is formed with biotin bound at one end and fluorescein bound at the other end. The labeled nucleic acid serves as an analyte in a two site immunometric assay. That is, a microparticle coated with anti-fluorescein antibodies captures the ligated product; after a wash step, a second biotin specific antibody labeled with alkaline phosphatase binds to the biotin yielding a fluorescent signal when incubated with a suitable substrate. The signal can be read with a fluorometer. This system is based on immunometric technology and requires specialized equipment

for detection of the target nucleic acid.

## SUMMARY OF INVENTION

This invention is based on a novel concept for a method for detecting specific DNA or RNA sequences. The present invention is defined by  
5 agglutination through the linking of microparticles with two distinct haptens, and alternatively, by linking microparticles to a capture zone on a lateral flow membrane or a filtration membrane with two distinct haptens.

The present invention is applicable to the detection of all nucleic acids and derivatives thereof. The present invention is useful to identify  
10 specific nucleic acid sequences associated with certain diseases or conditions, to screen amplicon detection for more efficient library screening and to detect the existence of amplification, but is not intended to be limited to these uses. This invention includes a detection system for any amplification-based diagnostic.

15 The present invention is a method for amplification and detection of a specific target nucleic acid comprising denaturing a double stranded nucleic acid and amplifying the target enzymatically with hapten derivatized complementary primers and dNTPs, forming a water-insoluble product of said amplified target nucleic acid and detecting the resulting product.

20 In one embodiment of this invention, nucleic acid detection by microparticle agglutination is accomplished by amplifying, by polymerase chain reaction (PCR), a specific sequence of the cytomegalovirus genome using biotin and digoxigenin primers.

In another embodiment of this invention, combining bifunctionality  
25 and microparticle technology is directed toward the creation of a visible line of detection on a lateral flow assay.

Other features and advantages of the present invention will become apparent from the following detailed description. taken in conjunction with the accompanying figures, that illustrate by way of example, the principles

of the instant invention.

### **BRIEF DESCRIPTION OF THE FIGURES**

5        Figure 1 illustrates the denatured nucleic acid target to be amplified with one primer 5' labeled with biotin and a second primer 5' labeled with digoxigenin.

Figure 2 shows the double strand PCR product resulting from amplification of the target nucleic acid sequence with the labeled primers.

Figure 3 illustrates large microparticles conjugated with streptavidin and small microparticles conjugated with anti-digoxigenin antibody.

10        Figure 4 illustrates the cross-linking and agglutination reaction in the presence of the amplified nucleic acid target sequence.

Figure 5 illustrates the assay results in the absence of an amplified target nucleic acid in the sample.

15        Figure 6A depicts the microscopic view of glass slide agglutination reactions; the negative control shows no visible agglutination.

Figure 6B depicts the microscopic view of glass slide agglutination reactions; visible agglutination indicative of positive results.

20        Figure 7 depicts naked eye views of agglutination reactions. Visible agglutination is indicative of positive results; the negative control shows no visible agglutination.



Figure 8A depicts the results of lateral flow membrane assays of bifunctionally labeled Human Immunodeficiency Virus (HIV) amplification product; a visible line of detection is not present in the negative control results.

5           Figure 8B depicts the results of lateral flow membrane assays of bifunctionally labeled Human Immunodeficiency Virus (HIV) amplification product; positive results are shown as a visible line of colored microparticles captured on the membrane.

10           Figure 9 depicts the results of lateral flow membrane assays of bifunctionally labeled amplification product. Positive results are shown as a visible line of colored microparticles captured on the membrane; a visible line of detection is not present in the negative control results.

#### **DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENT**

15           It is to be understood that both the foregoing general description and the following detailed description are exemplary and explanatory only and are not restrictive of the invention as claimed.

20           The present invention describes a method for the detection of nucleic acids. The present invention is embodied in a method of detecting an amplified target nucleic acid sequence that is present in a sample. The skilled artisan recognizes that assays for a broad range of target nucleic acid sequences present in a sample may be performed in accordance with the present invention. Samples may include biological samples derived from agriculture sources, bacterial and viral sources, and from human or other animal sources, as well as other samples such as waste or drinking water, agricultural products, processed foodstuff, air, etc. Examples include  
25           blood, stool, sputum, mucus, serum, urine, saliva, teardrop, a biopsy sample, an histological tissue sample, a tissue culture product, an agricultural

product, waste or drinking water, foodstuff, air, etc. The present invention is useful for the detection of nucleic acid sequences indicative of genetic defects or contagious diseases.

5 The following definitions will be helpful in understanding the specification and claims. The definitions provided herein should be borne in mind when these terms are used in the following examples and throughout the instant application.

As used herein, the term "target" nucleic acid molecule refers to the nucleic acid molecule that is to be amplified by the disclosed methodology. 10 The "target" molecule can be present in a purified, partially purified or unpurified state in the sample.

As used in this invention, the term "amplification" refers to a "template-dependent process" that results in an increase in the concentration of a nucleic acid sequence relative to its initial concentration. A "template-dependent process" is a process that involves the "template-dependent 15 extension" of a "primer" molecule. A "primer" molecule refers to a sequence of a nucleic acid which is complementary to a known portion of the target sequence and labeled with a hapten. A "template dependent extension" refers to nucleic acid synthesis of RNA or DNA wherein the sequence of the newly synthesized strand of nucleic acid is dictated by the 20 rules of complementary base pairing of the target nucleic acid and the primers.

The present invention relates to the amplification of one specific nucleic acid sequence, resulting in a nucleic acid duplex with discrete 25 hapten labels derivatized to each of the 5' ends or in the case of amplification by the ligase chain reaction, discrete haptens labeled on the 5' and 3' ends.

A second embodiment relates to the use of hybridization technology to create a bifunctionally labeled product. The probe may be singly labeled 30 and designed to generate bifunctional label when combined with primer that

is labeled or the probe itself may be bifunctionally labeled. This embodiment is useful under certain isothermal amplification conditions.

In the most preferred embodiment of the invention, a denatured target nucleic acid is amplified and 5' primer labeled, as illustrated in Figure 1.

5 The first primer consists of a short nucleic acid sequence complementary to a sequence on the target nucleic acid and a biotin label on the 5' end. The second primer consists of a nucleic acid sequence which is complementary to a second sequence on the target nucleic acid and a digoxigenin label on the 5' end. The results of amplifying the target nucleic acid with the primers  
10 in a template dependent extension reaction are shown in Figure 2.

In this embodiment, biotin and digoxigenin are incorporated onto the 5' ends of the resulting nucleic acid. Figure 3 shows large dyed  
microparticles conjugated with streptavidin and small dyed microparticles conjugated with anti-digoxigenin antibody, while Figure 4 illustrates the  
15 results of combining the amplified nucleic acid containing the biotin and digoxigenin labels with large dyed microparticles conjugated with streptavidin and small dyed microparticles conjugated with anti-digoxigenin antibody. The result is a linear and three dimensional cross-linking of the microparticles and labeled nucleic acid, thereby detecting the amplified  
20 nucleic acid sequence.

The results of a negative sample combined with the large dyed microparticles conjugated with streptavidin and small dyed microparticles conjugated with anti-digoxigenin antibody are shown in Figure 5. Since  
25 the target nucleic acid is not present in the sample, it cannot bind the primers required for the amplification process. The result is that some biotin derivatized primer will bind to the large dyed microparticles conjugated with streptavidin. Some digoxigenin derivatized primer will bind to the small dyed microparticles conjugated with anti-digoxigenin antibody. However, there is no linear or three dimensional cross-linking of the  
30 microparticles because the amplified nucleic acid is not present.

Accordingly, several objects and advantages of the invention described herein include: the rapid and accurate detection of amplified nucleic acid sequences with or without the use of hybridization and elaborate detection equipment; the assay disclosed herein is easy to perform, requiring little expertise in the art of molecular biology; the cost is significantly less than other methods known in the art; and the time frame for detecting an amplified sequence is cut drastically. As a research tool, this invention eliminates the need for preparing and running agarose gels to determine if a particular amplification method is successful. It may also be used in research in the screening of amplicon detection for the purpose of more efficiently screening libraries. As a diagnostic tool, the results are available a few minutes after amplification instead of hours or days. There is no danger from potential health hazards such as radioisotopes and ethidium bromide. The assay does not require special waste disposal procedures. The reagents employed are universal, meaning that the detection system in this invention can be used with any amplified nucleic acid containing bifunctional haptens specific for the receptor microparticles. The requirement for many washes in a immunometric or hybridization approach are eliminated.

Nucleic acid probe technology has developed rapidly in recent years as the scientific community has discovered its value in the detection of various diseases, organisms or genetic features. Amplification techniques provide the sensitivity to qualitatively determine the presence of even minute quantities of nucleic acid.

The invention disclosed herein relates generally to a detection method to qualitatively detect nucleic acid sequences which have been amplified from a selected target. This invention relies on the principles of bifunctional labeling of target nucleic acid during the amplification process, ligand-receptor binding, and microparticle technology for visible detection of amplified nucleic acid. The detection reaction is between the

bifunctional haptens on the amplicon and dyed microparticles conjugated with ligands specific for binding to each hapten.

The sensitivity and specificity of the process described herein does not necessarily rely on hybridization of the target nucleic acid with an oligonucleotide probe. The detection system is not fluorescent or enzyme based. The sensitivity and specificity of this invention depends on the incorporation of two distinct hapten labels during the amplification step or subsequent hybridization that generates a bifunctional label. If both haptens are not present the target nucleic acid will not be detected.

The process according to the present invention is suitable for the qualitative determination of all nucleic acid amplicons. The sensitivity and accuracy of this process is improved compared to the processes currently used by those skilled in the art. The invention offers the possibility of rapid and reliable qualitative determination of the presence of amplified target nucleic acids with simple reagents.

The following examples serve to explain and illustrate the present invention. Said examples are not to be construed as limiting of the invention in anyway. Various modifications are possible within the scope of the invention.

#### **Example 1 Derivatization and Labeling**

The present invention employs a variety of different enzymes to accomplish amplification of the target nucleic acid sequence. These enzymes are called polymerases and ligases. Polymerases are defined by their function of incorporating nucleoside triphosphates to extend a 3' hydroxyl terminus of a "primer molecule." As used herein, a "primer" is an oligonucleotide, that when hybridized to a target nucleic acid molecule, possesses a 3' hydroxyl terminus that can be extended by a polymerase and a hapten label at or near the 5' terminus. For a general discussion concerning polymerases, see Watson, J.D. *et al.*, (1987) *Molecular Biology*

of the Gene, 4th Ed., W.A. Benjamin, Inc., Menlo Park, CA. Examples of polymerases that can be used in accordance with the methods described herein include, but are not limited to, *E. coli* DNA polymerase I, the large proteolytic fragment of *E. coli* polymerase I, commonly known as

5 "Klenow" polymerase, Taq-polymerase, T7 polymerase, T4 polymerase, T5 polymerase and reverse transcriptase. The ligase chain reaction, described in European Patent No. 320308, uses 4 nucleic acid oligonucleotides that hybridize to the denatured target sequence, two on each strand (A and B), adjacent to one another and complementary to the two primers which

10 hybridize with the other strand (A' and B'). After hybridization, the two primers on each strand are joined with the enzyme, DNA ligase. In this embodiment of the invention, each oligonucleotide A and B' is derivatized with hapten (H<sub>1</sub>) on the 3' end and A' and B would need to be derivatized with hapten (H<sub>2</sub>) on the 5' end. The resulting double stranded product will

15 be labeled with (H<sub>1</sub>) and (H<sub>2</sub>).

In the present invention, polymerases with the capacity to continue the extension of a particular primer to produce a double stranded extension product with discreet hapten labels on each of the 5' ends are preferred. The general principles and conditions for amplification of nucleic acids using

20 polymerase chain reaction are quite well known, the details of which are provided in numerous references including United States Patent No. 4,683,195, United States Patent No. 4,683,202 and United States Patent No. 4,965,188, all to Mullis *et al.*, all of which are specifically incorporated herein. Thus, the details of PCR technology are not included herein.

25 However, in view of the teaching in the art and the specific teaching provided herein, a skilled artisan is able to practice the present invention by making only minor adjustments, as outlined herein, in order to accomplish the advantages of detection utilizing this system.

The oligonucleotide primers (P<sub>1</sub> and P<sub>2</sub>) used in this invention

30 contain at least one hapten, as label, which does not participate in the

priming reaction. The hapten is bound to at least one position of the nucleic acid primer. For the derivatization of nucleic acid primers, various methods can be employed (Maniatis *et al.* (1982) *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y.) The incorporation of the hapten can take place enzymatically, chemically or photochemically. The hapten can be derivatized directly to the 5' end of the primer or contain a bridge 1 to 30 atoms long. In the preferred embodiment, the bridge is linear. However, in a further embodiment, the bridge consists of a branched chain with a hapten molecule on at least one of the chain ends. By means of the presence of several hapten molecules on the ends of a branched chain, the detection sensitivity is increased. The preferred haptens for this invention are biotin ( $H_1$ ) and digoxigenin ( $H_2$ ), however, any other hapten which has available a receptor as specific binding agent is also suitable, for example, steroids, halogens and 2,4 dinitrophenyl.

The preferred method for utilization of the invention is a process for the detection of "target" nucleic acid sequence which is characterized in that the sample is treated, after denaturation, with two distinct labeled oligonucleotides primers ( $P_1$  and  $P_2$ ). The first ( $P_1$ ) is, in its sequence, complementary to a partial sequence of a strand of the target nucleic acid and is labeled with hapten ( $H_1$ ). An additional primer ( $P_2$ ) is complementary to another partial sequence of the target nucleic acid and labeled with a second hapten ( $H_2$ ). The mixture is treated with polymerase, preferably Taq-polymerase, deoxyribonucleotide and subsequently repeating, at least once, the cycle consisting of denaturation, hybridization and polymerization. A double stranded nucleic acid results which is labeled with two distinct haptens at each 5' end. The resulting labeled nucleic acid is reacted with conjugated microparticles of suitable color and diameter for detection. The first of which is conjugated with a receptor ( $R_1$ ) specific for binding at least one site on the first of the two distinct haptens ( $H_1$ ). The second is conjugated with a receptor ( $R_2$ ) specific for binding at least one

site on the second hapten ( $H_2$ ), the resulting microparticle bound amplicon is detected visually.

By way of example, cytomegalovirus is used as a PCR amplification target according to the PCR reaction and thermal cycling conditions disclosed by Gerdes (1993) *Transplantation Proceedings* 25:1411. PCR generated amplicons for bead detection utilize primers five-prime derivatized with either biotin or digoxigenin (Operon Technologies, Inc., Alameda, CA). Streptavidin coated 4.5  $\mu$ m beads are obtained from Perkin Elmer (Foster City, CA) and anti-digoxigenin beads are made using anti-dig Fab obtained from Boehringer Mannheim (Indianapolis, IN) and covalently bound to Dynal (Great Neck, NY) M-280 toxylactivated 2.8  $\mu$ m beads. The PCR amplicon is run on agarose gel, cut out and purified with GeneClean (Bio 101, Alameda, CA) to eliminate possible primer dimers artifacts.

## **Example 2    Microparticle Selection and Preparation**

The preferred microparticles utilized in this invention are composed of polymeric materials such as latex polyethylene, polypropylene, polymethylmethacrylate or polystyrene. However, a variety of other synthetic or natural materials may also be used in the preparation of the microparticles, for example, silicates, paramagnetic particles and colloidal gold. The usual form of microparticle possesses sulfate charge groups on their surface but these surfaces can be modified by the introduction of functional groups such as hydroxyl, carboxyl, amine and carboxylate groups. The functional groups are used to bind a wide variety of ligands and receptors to the microparticles. These groups are selected on the basis of their ability to facilitate binding with the selected member of the ligand-receptor pair, either by covalent binding or adsorption. The preferred method of attachment of the receptor to the microparticles is covalent binding.

The size of the microparticles used in this invention is selected to



optimize the binding and detection of the labeled amplicons. Microparticles are available in a size range of 0.02-100  $\mu\text{m}$  in diameter. The preferred diameter for this embodiment of the invention is the range of 0.1-1.0  $\mu\text{m}$ , specifically not excluding the use of either larger or smaller microparticles as appropriately determined. The microparticles are activated with a suitable receptor for binding to the target ligand. The preferred microparticle in the present invention is composed of latex containing a colored dye.

In the present invention, microparticle bound receptors are specific for discrete haptens located on the ends of amplified nucleic acid sequences. The receptors must be capable of binding to their specific binding partner (hapten) and, further changing the derivatized haptens from the preferred biotin and digoxigenin necessitates a change in the receptors. Conjugation of the receptors to the microparticle is accomplished by covalent binding or, in appropriate cases, by adsorption of the receptor onto the surface of the microparticle. Techniques for the adsorption or covalent binding of receptors to microparticles are well known in the art and require no further explanation.

In order to prepare the anti-digoxigenin coated microparticles, 0.25-1.0 mg/ml of anti-digoxigenin Fab is incubated with a suspension containing a final concentration of 1.0% microparticles/ml. The microparticles and digoxigenin Fab are allowed to react for 15 minutes prior to treatment with activating agent for covalent binding. The microparticles are treated with EDAC (1-ethyl-3-(3-dimethylaminopropyl)carbodiimide) at a final concentration of 0-2.5 mM. The Fab and microparticles are then mixed and incubated at room temperature for one hour. Unbound Fab is removed by successive washes and resuspended in storage buffer. The sensitivity of the coated microparticles is determined by: 1) agglutination reaction between secondary anti-body to digoxigenin Fab; 2) lateral flow assay; and 3) filtration assay. Inhibition of the reaction is demonstrated by

pre-incubation with digoxigenin Fab and secondary anti-body to digoxigenin Fab, prior to adding coated microparticles.

5 The microparticles are 0.3-10.0  $\mu\text{m}$  in diameter. Three different carboxyl group densities and three different colors are evaluated for reactivity. The reactivity is evaluated by agglutination reactions, lateral flow membrane assay and filtration assay.

10 Agglutination reactions consist of adding 10  $\mu\text{l}$  of each reactant to 10  $\mu\text{l}$  of reaction buffer on a glass slide. The reaction is mixed gently and the rate of agglutination determined. Positive reactions show visible agglutination within 2 minutes and are rated on a 0 to +4 scale. Negative controls do not have visible agglutination within the specified time frame. Microparticles coated with non-specific protein or leaving the "bridging component" out of the reaction are used as negative controls.

15 Lateral flow membrane assays are performed on Biodyne C nylon membranes (Pall BioSupport Division, Port Washington, NY) spotted with 1.0  $\mu\text{l}$  streptavidin at concentrations between 0.0 and 1.0 mg/ml. Next, 25-75  $\mu\text{l}$  of colored, biotinylated, coated beads, at dilutions of 0.001-1.0% microparticles/ml are allowed to wick up the membrane. A positive reaction results in a colored spot where the capture material is applied. 20 Membranes spotted with 1.0  $\mu\text{l}$  of buffer serve as negative controls. The membranes show background when soaked in a suitable blocking reagent, such as bovine serum albumin (BSA), "Tween-20" or Casein.

25 The filtration assays are accomplished by adding 10  $\mu\text{l}$  of each reactant to 10  $\mu\text{l}$  of reaction buffer in an ELISA well or microfuge tube, or spotting the reaction on a glass slide. The reaction proceeds for 2-10 minutes at room temperature. A 1.0-5.0  $\mu\text{l}$  sample is removed from the reaction and spotted on a membrane filter, said filter being of a suitable pore size such that non-agglutinated microparticles pass through, but agglutinated microparticles cannot. The filter is then washed 2 times with 30 phosphate buffered saline (PBS). Positive results are visible as colored

"spots" on the filter, while negative results are not visible. That is, no colored "spots" are produced on the filters.

### **Example 3 Target Nucleic Acid Amplicon Detection**

5 The present invention includes the processes for the qualitative determination of specific target nucleic acid amplicons which have been bifunctionally labeled during amplification by incubating the hapten labeled target nucleic acid with at least two receptors, ( $R_1$ ) and ( $R_2$ ), whereby ( $R_1$ ) and ( $R_2$ ) are incapable of binding to each other and ( $R_1$ ) and ( $R_2$ ) are bound to dyed microparticles. The haptens disclosed herein, ( $H_1$ ) and ( $H_2$ ), are  
10 incapable of binding to each other. In the instant invention, ( $R_1$ ) specifically binds to at least one site on ( $H_1$ ) and ( $R_2$ ) specifically binds to at least one site on ( $H_2$ ). Detection of the target nucleic acid amplicon is accomplished in several ways as described below. Preferred receptors are streptavidin ( $R_1$ ) and anti-digoxigenin antibody ( $R_2$ ).

15 In a first embodiment, ( $R_1$ ) and ( $R_2$ ) conjugated dyed microparticles are added to a portion of the amplicon reaction mixture on a surface that does not bind any of the components of the assay, for example, a glass slide. The result, a linear and three dimensional cross-linkage of the amplicon. The resulting agglutination reaction is visible to the naked eye or by  
20 observation with an amplifying device such as a microscope. The reaction takes place in a matter of minutes. The specificity and sensitivity of the assay are determined by the primers. Agglutination can only take place if the target nucleic acid is present in the amplification mixture to form cross-links between the beads. If the target nucleic acid is not present, the dyed  
25 microparticles remain monodispersed and agglutination is not visible.

Agglutination reactions consist of depositing 10  $\mu$ l of 1% anti-digoxigenin coated microparticles, 10  $\mu$ l of 1% streptavidin coated microparticles, 10  $\mu$ l of reaction buffer and 10  $\mu$ l of diluted HIV amplicon or buffer on a glass slide. The reaction is gently mixed and the rate of

agglutination determined over a period of 2 minutes. Agglutination is visible in the reaction mix containing HIV amplicon. The negative control, containing buffer in place of amplicon, does not show agglutination. An additional negative control uses microparticles coated with non-specific protein. The results of such reactions are shown in Figures 6 and 7.

Another embodiment includes a method of detection relying upon filtration of the agglutination reaction through a porous membrane, said membrane having pores of a size such that unagglutinated microparticles flow through the membrane but agglutinated microparticles are retained on the filter as a visible "spot."

Filtration assays are performed by placing 10  $\mu$ l of 1% anti-digoxigenin coated microparticles, 10  $\mu$ l of 1 % streptavidin coated microparticles, 10  $\mu$ l of reaction buffer and 10  $\mu$ l of diluted FHV amplicon or buffer in an ELISA well or microfuge tube, or spotting the reaction components on a glass slide. The reaction proceeds for 2-10 minutes at room temperature. A 1.0-5.0  $\mu$ l sample is removed from the reaction and spotted on a membrane filter, said filter being of a suitable pore size such that nonagglutinated microparticles pass through but agglutinated microparticles cannot. The filter is then washed 2 times with PBS. Positive results are visible as colored "spots" on the filter, while negative results are not visible.

An alternative embodiment is to anchor one of the receptors ( $R_1$ ) to a solid phase membrane. This group of receptors serves to capture the target sequence. The amplified nucleic acid is mixed with the second receptor ( $R_2$ ) that is conjugated to a dyed microparticle. If the target nucleic acid is present in the reaction mix, ( $R_2$ ) binds to ( $H_2$ ) on the amplified nucleic acid. This mixture is then deposited on the membrane containing ( $R_1$ ) and allowed to interact. If the target is present, the anchored receptor ( $R_1$ ) interacts with hapten ( $H_1$ ) to capture the amplified nucleic acid bound to the dyed microparticle through the interaction of ( $H_2$ ) and ( $R_2$ ). The result is a

line of dyed particles visible on the membrane. If the target is not present, the dyed particle is not captured on the membrane, and thus, is not visible.

**Example 4 Lateral Flow Chromatography of Bifunctionally Labeled Human Immunodeficiency Virus (HIV) Amplification Product**

In the instant set of experiments, the primers, SK 38 and SK 39, are obtained from Operon, of Alameda CA, and the derivatization methodology employed is outlined previously, in Example 1. The forward primer, SK 38, carries a 5' digoxigenin (dig) label. The reverse primer, SK 39, carries a biotin 5' label. The sequences of these primers are shown below:

**SK 38 dig primer**

5'-dig ATA ATC CAC CTA TCC CAG TAG GAG AAA T-3' SEQ ID NO:1

**SK 39 biotin primer**

5'-biotin TT TGG TCC TTG TCT TAT GTC CAG AAT GC-3 SEQ ID NO:2

The sequence target for amplification of HIV, positions 1090 to 1205, is as follows:

5' ATAATCCACCTATCCAGTAGGAGAAATTTATAAAAGATGGATAATCCTGGGATTAAATAAAAT  
AGTAAGAATGTATAGCCCTACCAGCATTCTGGACATAAGACAAGGACCAAA SEQ ID NO:3

Membrane used for this procedure is nitrocellulose, purchased from Millipore Corporation of Bedford, MA. A stripe of streptavidin at a concentration of 1 mg/ml is applied at a rate of 1 µl/cm via a linear reagent striper (IVEK Corporation, No. Springfield, VT) 1 cm from the bottom edge of the membrane. After application of the streptavidin, the membrane is allowed to dry and then blocked for non-specific binding by 0.5% casein in 100 mM Tris, pH 7.4. The membranes are washed twice with water

(ddH<sub>2</sub>O) and allowed to dry. The microparticles used are prepared as outlined previously, in Example 2.

The target is amplified by PCR using 200-1000 mM primer concentration, GeneAmp EZ rTth RNA PCR kit (Perkin Elmer Corp., Alameda, CA) and 10<sup>3</sup>-10<sup>6</sup> copies/ml of the target sequence. Thirty to forty PCR cycles, each cycle being 60°C for 15 minutes, 95°C for 15 seconds, and 55°C for 60 seconds, are run. The resulting amplicon is bifunctionally labeled during the PCR reaction with biotin and digoxigenin.

The amplicon (5 µl) is added to 5 µl of 0.1% anti-digoxigenin coated microparticles and 40 µl of water (ddH<sub>2</sub>O), then applied (50 µl) to the previously striped membrane. The amplicon bound to the anti-digoxigenin microparticles wicks through the membrane to the streptavidin line and is captured by the interaction of biotin and streptavidin. The result is a visible line of colored microparticles.

In the negative control, the procedure is performed as described above, but without the addition of the target sequence. Without the presence of the target sequence in the amplification reaction, the bifunctionally labeled amplicon is not generated and the visible line of detection is not present. The results of such experiments are shown in Figures 8 and 9.

**Example 5 Lateral Flow Chromatography of Bifunctionally Labeled Cytomegalovirus (CMV) Amplification Product**

The primers used in this set of experiments. MIEAN and MIEBN, are from Operon of Alameda, CA. The derivatization methodology employed is outlined previously, in Example 1. The forward primer, MIEAN, carries a 5' digoxigenin label. The reverse primer, MIEBN, carries a biotin 5' label. The sequences of these primers are shown below:

## MIEAN

5' digGCATTGAGGAGATCTGCATGAAGG

SEQ ID NO:4

## MIEBN

5' biotinATTACTGAGGACAGAGGGATAGTC

SEQ ID NO:5

5           The sequence target for amplification of CMV is nucleotide 2758-3060.

          Membrane used in this procedure is nitrocellulose, purchased from Millipore Corporation of Bedford, MA. A stripe of streptavidin at a concentration of 1 mg/ml is applied at a rate of 1  $\mu$ l/cm via a linear reagent  
10           striper (IVEK Corporation, No. Springfield, VT) 1 cm from the bottom edge of the membrane. After application of the streptavidin, the membrane is allowed to dry and then blocked for non-specific binding by 0.5% casein in 100 mM Tris, pH 7.4. The membranes are washed twice with water (ddH<sub>2</sub>O) and allowed to dry. The microparticles used are prepared as  
15           outlined previously, in Example 2.

          The target is amplified by PCR using 200-1000 mM primer concentration, GeneAmp EZ rTth RNA PCR kit (Perkin Elmer Corp., Alameda, CA) and 10<sup>3</sup>-10<sup>6</sup> copies/ml of the target sequence. Thirty to forty  
20           PCR cycles, each cycle being 60°C for 15 minutes, 95°C for 15 seconds, and 55°C for 60 seconds, are run. The resulting amplicon is bifunctionally labeled during the PCR reaction with biotin and digoxigenin.

          The amplicon (5  $\mu$ l) is added to 5  $\mu$ l of 0.1% anti-digoxigenin coated microparticles and 40  $\mu$ l of water (ddH<sub>2</sub>O), then applied (50  $\mu$ l) to the previously striped membrane. The amplicon bound to the anti-digoxigenin  
25           microparticles wicks through the membrane to the streptavidin line and is captured by the interaction of biotin and streptavidin. The result is a visible line of colored microparticles.

          In the negative control, the procedure is performed as described

above, but without the addition of the target sequence. Without the presence of the target sequence in the amplification reaction, the bifunctionally labeled amplicon is not generated and the visible line of detection is not present.

5       **Example 6   Agglutination of HIV Amplification Product Using One Labeled Primer and One Unlabeled Primer, Followed by Hybridization with a Second Labeled Probe in Order to Create a Bifunctionally Labeled Amplicon**

10       The primers, SK 38 and SK 39, and the sequence target for amplification of HIV used in this example are identical to those described in Example 4 (SEQ ID NOS: 1-3). Furthermore, the derivatization methodology is outlined previously, in Example 1. The sequence of the 61mer probe employed in this example is as follows:

61mer probe

15       5'-(Biotin)GCT GGT AGG GCT ATA CAT TCT TAC TAT TTT ATT  
TAA TCC CAG GAT TAT CCA TCT TTT ATA A-3'                      SEQ ID NO:6

20       The target is amplified by PCR using 200-1000 mM primer concentration, GeneAmp EZ rTth RNA PCR kit (Perkin Elmer Corp., Alameda, CA) and  $10^3$ - $10^6$  copies/ml of the target sequence. Thirty to forty PCR cycles, each cycle being 60°C for 15 minutes, 95°C for 15 seconds, and 55°C for 60 seconds, are run. The resulting amplicon is labeled only with biotin during the PCR reaction.

25       Following amplification, 5 µl of SK 38 Dig---SK 39 single labeled product is added to 5 µl of 25 µM (125 pmol) SK 39 labeled with biotin. The mixture is heated to 95°C for 1 min, then cooled to 55°C for 1 min. Alternatively, 5 µl of 1:2 dilution in water of the SK 38 Dig----SK 39 single labeled product is added to 5 µl of 2.5 µM HIV 61mer biotin labeled probe. The mixture is heated to 95°C for 1 min, then cooled to 60°C for 1 min. The



hybridization step constructs a product that is bifunctionally labeled.

The microparticles are prepared as outlined previously, in Example 2. Agglutination on a glass slide is accomplished by adding 5 µl of either hybridized product to 5 µl anti-dig coated microparticles (0.2%) and 5 µl streptavidin coated microparticles (0.2%). The results of a hybridization test containing a target sequence are a visible agglutination of the microparticles. Samples that do not contain a target sequence do not agglutinate and samples that are not bifunctionally labeled with the hybridization step also do not agglutinate.

10 **Example 7 Inhibition assay: Loss of Visible Signal on Lateral Flow Membrane**

The specific probe and target employed in the instant example have been designed by ID Biomedical Corporation for use in detecting *Mycobacterium tuberculosis*. The probe is a chimeric construct containing both DNA and RNA sequences with labels on the 5' (fam) and the 3' (biotin) ends of the DNA portion of the probe. The binding of the probe to a single strand of target generates double stranded nucleic acid which is cleaved with RNase H, thus, eliminating the bifunctionality of the probe. The sequence of the probe is described below:

20 **FARK2S3B probe**

5' - fam AAA GAT GT agag GGT ACA GA-3'biotin SEQ ID NO:7  
(lower case indicates deoxyribonucleoside bases)

The sequence of the target is described below:

ARK2-T synthetic target

25 5' - AAT CTG TAC CCT CTA CAT CTT TAA-3' SEQ ID NO:8

The reaction is completed following the protocol provided by ID Biomedical Corporation.

Membrane used for this procedure is nitrocellulose, purchased from Millipore Corporation of Bedford, MA. A stripe of streptavidin at a  
5 concentration of 1 mg/ml is applied at a rate of 1  $\mu$ l/cm via a linear reagent  
striper (IVEK Corporation, No. Springfield, VT) 1 cm from the bottom edge  
of the membrane. After application of the streptavidin, the membrane is  
allowed to dry and then blocked for nonspecific binding by 0.5% casein in  
100 mM Tris, pH 7.4. The membranes are washed twice with water  
10 (ddH<sub>2</sub>O) and allowed to dry. The microparticles used are prepared as  
outlined previously, in Example 2, replacing anti-digoxigenin Fab with anti-  
fam monoclonal IgG.

The reaction product (10  $\mu$ l) is added to 5  $\mu$ l of 0.1% anti-fam coated  
microparticles (0.1%) and 35  $\mu$ l of water (ddH<sub>2</sub>O), then applied (50  $\mu$ l) to  
15 the previously striped membrane. The binding of the probe to the target  
followed by cleavage of the probe by RNase H. results in loss of the  
bifunctionality of the probe. When the target is present, the absence of a  
visible line on the membrane exists. When the target is not present, the  
bifunctionally labeled probe is able to bind the anti-fam coated  
20 microparticles and the streptavidin bound to the membrane, resulting in a  
visible line.

**Example 8 Isothermal Amplification Approach to Detection with  
Bifunctionally Labeled Amplified Target Sequence**

The instant strand displacement assay (SDA) is an example of an  
25 isothermal amplification that can be detected by using microparticles and  
bifunctionally labeled product. SDA technology is described in United  
States Patent No. 5,455,166 to Becton Dickinson and Company,  
specifically incorporated herein.

This set of experiments is conducted with composite extension

primers that are labeled with biotin, fam or digoxigenin. Bumper primers are the same sequence as provided by Becton Dickinson and Company (Franklin Lakes, New Jersey). The sequences of the target, the bumper primer and the composite extension primer are as follows:

5 Bumper primers:

B1: 5'-CGATCGAGCAAGCCA SEQ ID NO:9

B2: 5'-CGAGCCGCTCGCTGA SEQ ID NO:10

Composite extension primers:

S1: 5'-fam/dig-ACCGCATCGAATGCATGTCTCGGGTAAGGCGTACTCGA SEQ ID NO:11

10 S2: 5'-biotin-CGATTCCGCTCCAGACTTCTCGGGTGTACTGAGATCCC SEQ ID NO:12

Target sequence:

5'TGGACCCGCCAACAAGAAGGCGTACTCGACCTGAAAGACGTTATCCACCAT  
ACGGATAGGGGATCTCAGTACACATCGATCCGGTTCAGCG SEQ ID NO:13

15 The reaction is set up per the thermophilic Strand Displacement Amplification (tSDA) protocol developed by Becton Dickinson and Co. The target organism is *Mycobacterium tuberculosis*. For pilot studies, an artificial target template consisting of the 91nt sequence of the *M. tuberculosis* genome, defined by the Becton Dickinson outer (bumper) primers, is used. Amplification conditions used are identical to those  
20 used by Becton Dickinson for tSDA.

Membrane used for this procedure is nitrocellulose, purchased from Millipore Corporation of Bedford, MA. A stripe of streptavidin at a concentration of 1 mg/ml is applied at a rate of 1 µl/cm via a linear reagent strip (IVEK Corporation, No. Springfield, VT) 1 cm from the bottom edge  
25 of the membrane. After application of the streptavidin, the membrane is

allowed to dry and then blocked for non-specific binding by 0.5% casein in 100 mM Tris, pH 7.4. The membranes are washed twice with water (ddH<sub>2</sub>O) and allowed to dry. Next, 3 µl of anti-S1 (complementary to S1 without the biotin label) and/or S2 primer (complementary to S2 without the dig or fam label) is spotted onto a second membrane. This membrane is sandwiched onto the first membrane in order to capture free primers that compete with the product for the microparticles or streptavidin capture zone. The microparticles used are prepared as outlined previously, in Example 2, with either anti-digoxigenin Fab or anti-fam monoclonal IgG. The microparticles are diluted 1:2 with a 35% sucrose solution and 3 µl applied directly to the membrane and dried.

The reaction product (10 µl) is added to 45 µl SDA buffer, then applied (50 µl) to the previously striped membrane. Application of the sample requires the bifunctionally labeled product and the competing primers to pass through the anti-primer coated membrane and the dried microparticles. When the target is present, there is a visible line on the membrane. When the target is not present, there is absence of a visible line.

Tests kits for the detection of target nucleic acid sequence determination that include hapten labeled primers, enzymes or ligases, nucleosides and receptor bound dyed microparticles are envisioned. Depending upon the detection method employed in each test kit, also included in the appropriate kit will be microscope slides, microporous membrane/filtration apparatus or microporous membrane/dip stick apparatus.

The instant invention provides a rapid, simple and accurate method of detecting amplified target nucleic acid sequences with a visual signal. The sensitivity and specificity of the assay are not based on hybridization but on the bifunctional labeling of the target during the amplification process. The method is as sensitive as current methodologies utilizing enzymes and fluoreceins. The method does not require costly and

sophisticated equipment or specially trained personnel and it does not pose a health hazard.

5 While the above description contains many specificities, these specificities should not be construed as limitations on the scope of the invention, but rather an exemplification of the preferred embodiment thereof. That is to say, the foregoing description of the invention is exemplary for purposes of illustration and explanation. Without departing from the spirit and scope of this invention, one skilled in the art can make various changes and modifications to the invention to adapt it to  
10 various usages and conditions. For example, other variations may include amplifying several target samples in the same reaction mixture, amplifying with ligase chain reaction, etc. As such, these changes and modifications are properly, equitably, and intended to be within the full range of equivalence of the following claims. Thus the scope of the  
15 invention should be determined by the appended claims and their legal equivalents, rather than by the examples given.

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT: John C. Gerdes
- 5 (ii) TITLE OF INVENTION: DETECTION OF AMPLIFIED  
NUCLEIC ACID SEQUENCES  
USING BIFUNCTIONAL  
HAPTENIZATION AND DYED  
MICROPARTICLES
- (iii) NUMBER OF SEQUENCES: 13
- 10 (iv) CORRESPONDENCE ADDRESS:  
(A) ADDRESSEE: Julie L. Bernard, Attorney  
at Law  
(B) STREET: 9000 E. Inspiration Drive  
(C) CITY: Parker  
15 (D) STATE: Colorado  
(E) COUNTRY: USA  
(F) ZIP: 80134-8535
- (v) COMPUTER READABLE FORM:  
20 (A) MEDIUM TYPE: Diskette, 3.5 inch  
(B) COMPUTER: IBM PC Compatible  
(C) OPERATING SYSTEM: MS-DOS  
(D) SOFTWARE: WordPerfect 6.0
- (vi) CURRENT APPLICATION DATA:  
25 (A) APPLICATION NUMBER:  
(B) FILING DATE: 12 July 1996  
(C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:  
(A) APPLICATION NUMBER: 60/002245  
(B) FILING DATE: 13 July 1995
- 30 (viii) ATTORNEY/AGENT INFORMATION:  
(A) NAME: Julie L. Bernard  
(B) REGISTRATION NUMBER: 36,450  
(C) REFERENCE/DOCKET NUMBER: IAD-2PCT
- 35 (ix) TELECOMMUNICATION INFORMATION:  
(A) TELEPHONE: 303 841 7472  
(B) TELEFAX: 303 840 1567

- (2) INFORMATION FOR SEQ ID NO:1:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 27 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
ATAATCCACC TATCCCAGTA GGAGAAA 27
- (2) INFORMATION FOR SEQ ID NO:2:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 28 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:2:  
TTTGGTCCTT GTCTTATGTC CAGAATGC 28
- (2) INFORMATION FOR SEQ ID NO:3:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 115 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:3:  
ATAATCCACC TATCCCAGTA GGAGAAATTT ATAAAAGATG  
GATAATCCTG GGATTAAATA AAATAGTAAG AATGTATAGC  
CCTACCAGCA TTCTGGACAT AAGACAAGGA CCAA 115
- (2) INFORMATION FOR SEQ ID NO:4:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 24 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:4:  
GCATTGAGGA GATCTGCATG AAGG 24
- (2) INFORMATION FOR SEQ ID NO:5:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 24 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:5:  
ATTACTGAGG ACAGAGGGAT AGTC 24

- 5 (2) INFORMATION FOR SEQ ID NO:6:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 28 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:6:  
GCTGGTAGGG CTATACATTC TTACTATTTT ATTTAATCCC  
AGGATTATCC ATCTTTTATA A 61
- 10 (2) INFORMATION FOR SEQ ID NO:7:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 20 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
15 (D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:7:  
AAAGATGTag agGGTACAGA 20
- 20 (2) INFORMATION FOR SEQ ID NO:8:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 24 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
25 (ix) SEQUENCE DESCRIPTION: SEQ ID NO:8:  
AATCTGTACC CTCTACATCT TTAA 24
- 30 (2) INFORMATION FOR SEQ ID NO:9:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 15 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
35 (ix) SEQUENCE DESCRIPTION: SEQ ID NO:9:  
CGATCGAGCA AGCCA 15
- 40 (2) INFORMATION FOR SEQ ID NO:10:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 15 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
45 (ix) SEQUENCE DESCRIPTION: SEQ ID NO:10:  
CGAGCCGCTC GCTGA 15



- 5 (2) INFORMATION FOR SEQ ID NO:11:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 40 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:11:  
ACCGCATCGA ATGCATGTCT CGGGTAAGGC GTACTCGACC40
- 10 (2) INFORMATION FOR SEQ ID NO:12:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 40 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
15 (D) TOPOLOGY: linear  
(ix) SEQUENCE DESCRIPTION: SEQ ID NO:12:  
CGATTCCGCT CCAGACTTCT CGGGTGTACT GAGATCCCCT40
- 20 (2) INFORMATION FOR SEQ ID NO:13:  
(i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 91 bases  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear  
25 (ix) SEQUENCE DESCRIPTION: SEQ ID NO:13:  
TGGACCCGCC AACAAGAAGG CGTACTCGAC CTGAAAGACG  
TTATCCACCA TACGGATAGG GGATCTCAGT ACACATCGAT  
CCGGTTCAGC G

**We Claim:**

1. A method for the detection of amplified target nucleic acid sequences which comprises:
  - 5 a) denaturing a multiple stranded nucleic acid to produce a single stranded nucleic acid or beginning with a single stranded nucleic acid;
  - b) amplifying the target nucleic acid sequence with hapten derivatized complimentary primers and dNTPs to produce bifunctionally labelled specific target nucleic acid amplicons  
10 specific for receptor conjugated dyed microparticles;
  - c) contacting receptor bound dyed microparticles and labelled target nucleic acid sequence wherein agglutination product of the amplified target nucleic acid is formed; and
  - d) detecting the resulting agglutination product.
- 15 2. The method of claim 1 wherein the primers are derivatized with biotin.
3. The method of claim 1 wherein the primers are derivatized with biotin and digoxigenin.
4. The method of claim 1 wherein the primers are derivatized with biotin and fam.  
20
5. The method of claim 1 wherein the primers are derivatized with biotin, fam or digoxigenin.
6. The method of claim 1 wherein the primers are derivatized with haptens selected from a group consisting of steroids, halogens and 2,4  
25 nitrophenyl.

7. The method of claim 1 wherein the amplification target is any specific nucleic acid sequence.
8. The method of claim 3 wherein the amplification target is Cytomegalovirus.
- 5 9. The method of claim 3 wherein the amplification target is Human Immunodeficiency Virus.
- 10 10. The method of claim 5 wherein the amplification target is *Mycobacterium tuberculosis*.
11. The method of claim 1 wherein the target nucleic acid is amplified isothermally.
12. The method of claim 1 wherein the dyed microparticles are covalently bound to hapten specific receptors.
13. The method of claim 1 wherein the amplified target nucleic acid is contacted with receptor bound dyed microparticles during amplification.
- 15 14. The method of claim 1 wherein the dyed microparticles are attached to the receptors via adsorption to the microparticles' surface.
15. The method of claim 12 wherein the bound receptors are streptavidin and anti-digoxigenin antibody.
- 20 16. The method of claim 12 wherein the bound receptors are streptavidin and anti-fam antibody.

17. The method claim 1 wherein the microparticles are 0.02-10.0 micrometers in diameter.
18. The method of claim 1 wherein the microparticles are 0.01-1.0 micrometers in diameter.
- 5 19. The method of claim 1 wherein the microparticles are composed of an insoluble polymer material.
20. The microparticles of claim 19 wherein the insoluble polymer material is latex.
21. The microparticles of claim 19 wherein the insoluble polymer material is selected from a group consisting of polyethylene,  
10 polypropylene, polymethylmethacrylate and polystyrene.
22. The method of claim 1 wherein the microparticles are composed of colloidal gold.
23. The method of claim 1 wherein said detection comprises  
15 determination of a presence or an absence of agglutination product, wherein said presence is indicative of amplified target nucleic acid.
24. The method of claim 1 wherein said agglutination product is detected using a secondary detection probe capable of binding to receptor molecules.
- 20 25. The method of claim 24 wherein the secondary detection probe is an anti-species antibody.

26. The detection of claim 23 wherein said determination is made by a detection method selected from a group consisting of agglutination, lateral flow membrane assay and filtration assay.
27. A kit for the amplification and detection of a target nucleic acid  
5 consisting essentially of:
- a) target specific hapten labeled primers;
  - b) thermostable polymerase;
  - c) dNTPs; and
  - d) receptor bound dyed microparticles.
- 10 28. The kit of claim 27 further consisting of agglutination product detection apparatus.
29. The agglutination product detection apparatus of claim 28 wherein said apparatus is a microporous membrane.
- 15 30. The microporous membrane of claim 29 wherein said membrane is nitrocellulose or nylon.
31. The microporous membrane of claim 30 wherein a receptor is immobilized thereon.

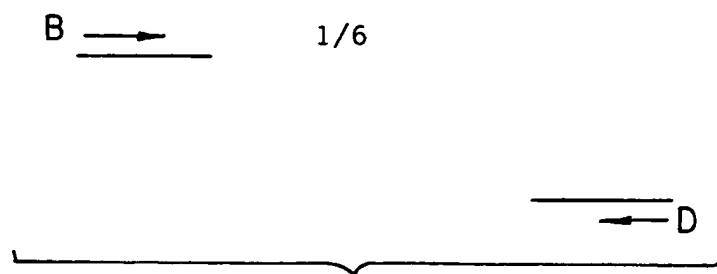


FIG. 1

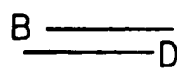


FIG. 2

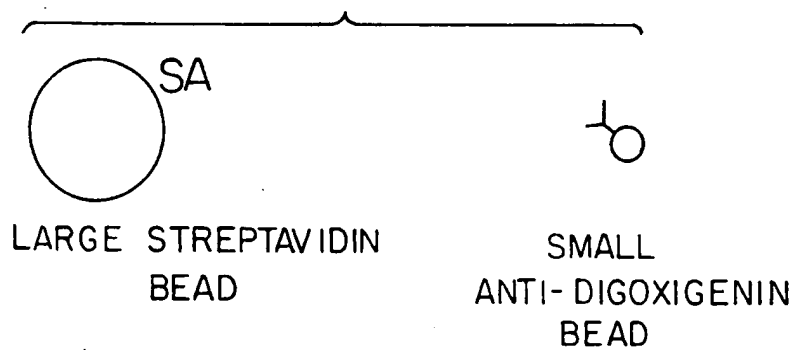


FIG. 3

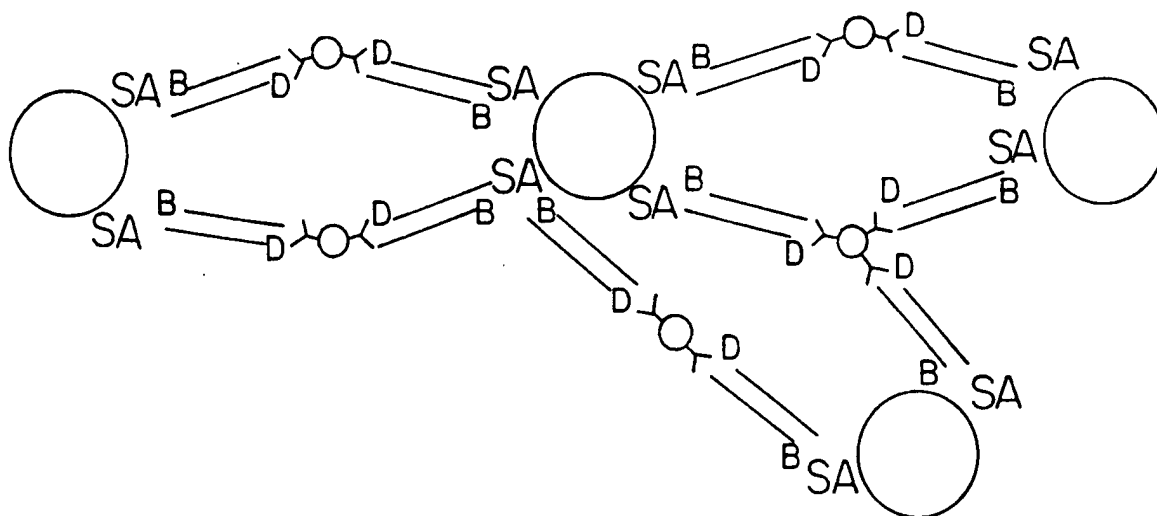


FIG. 4

2/6

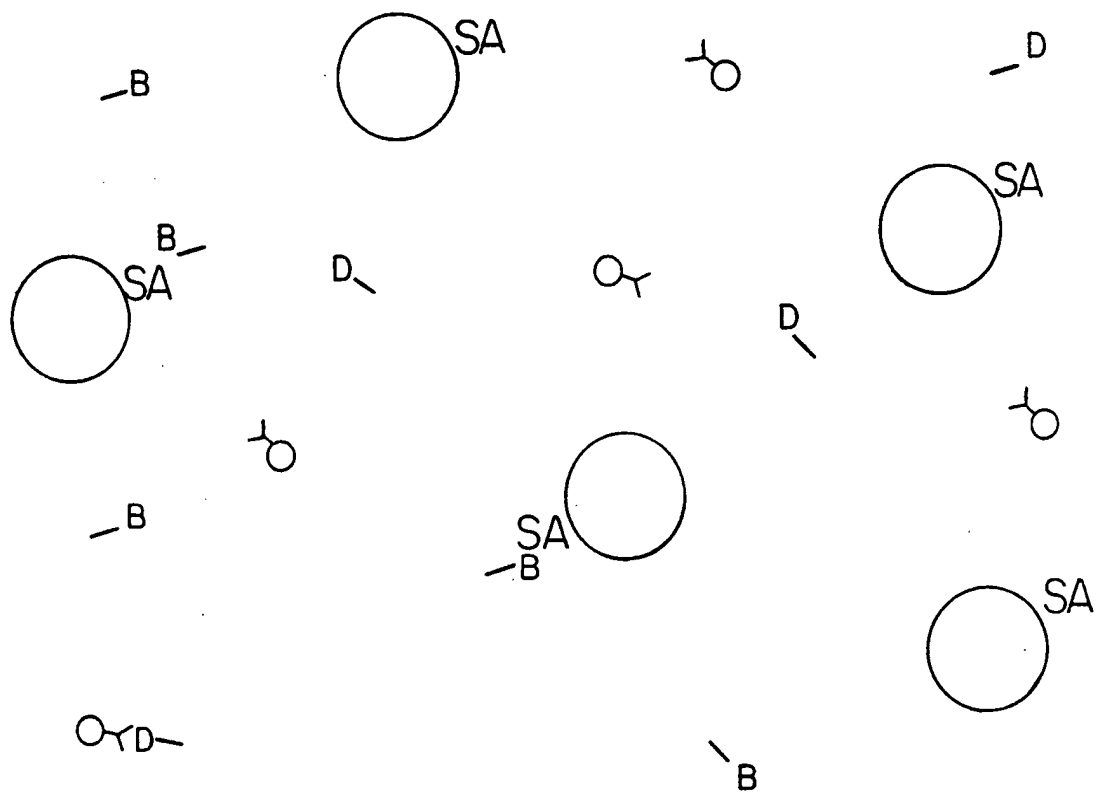


FIG. 5

3/6

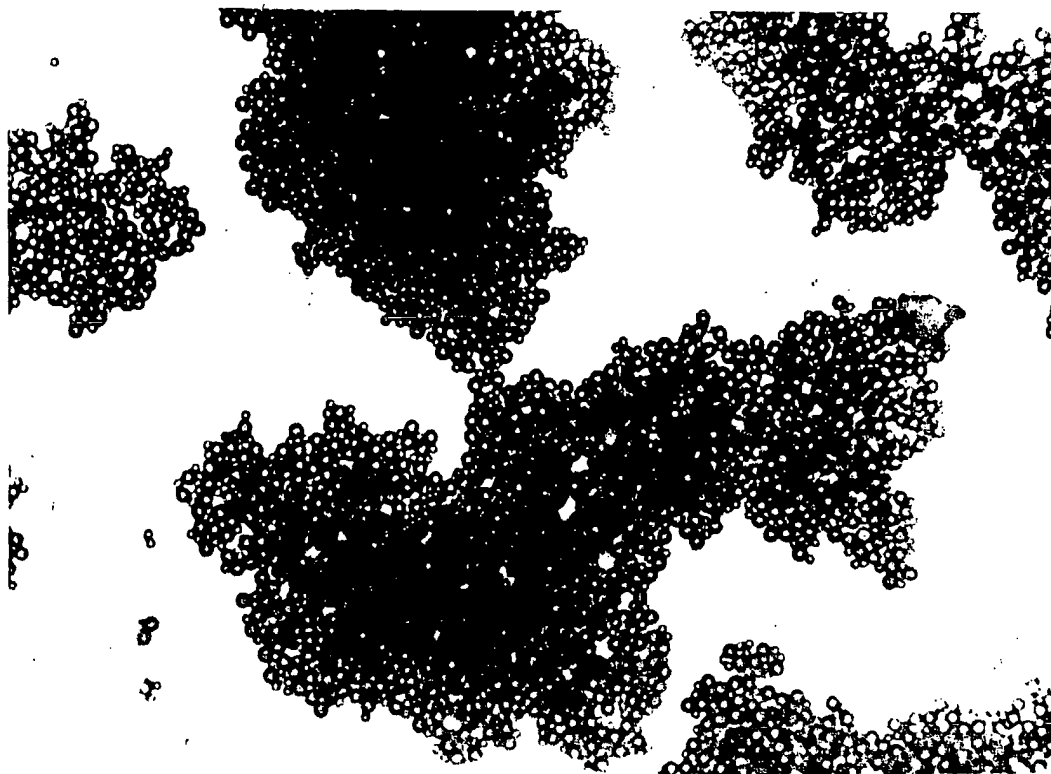


FIG. 6A

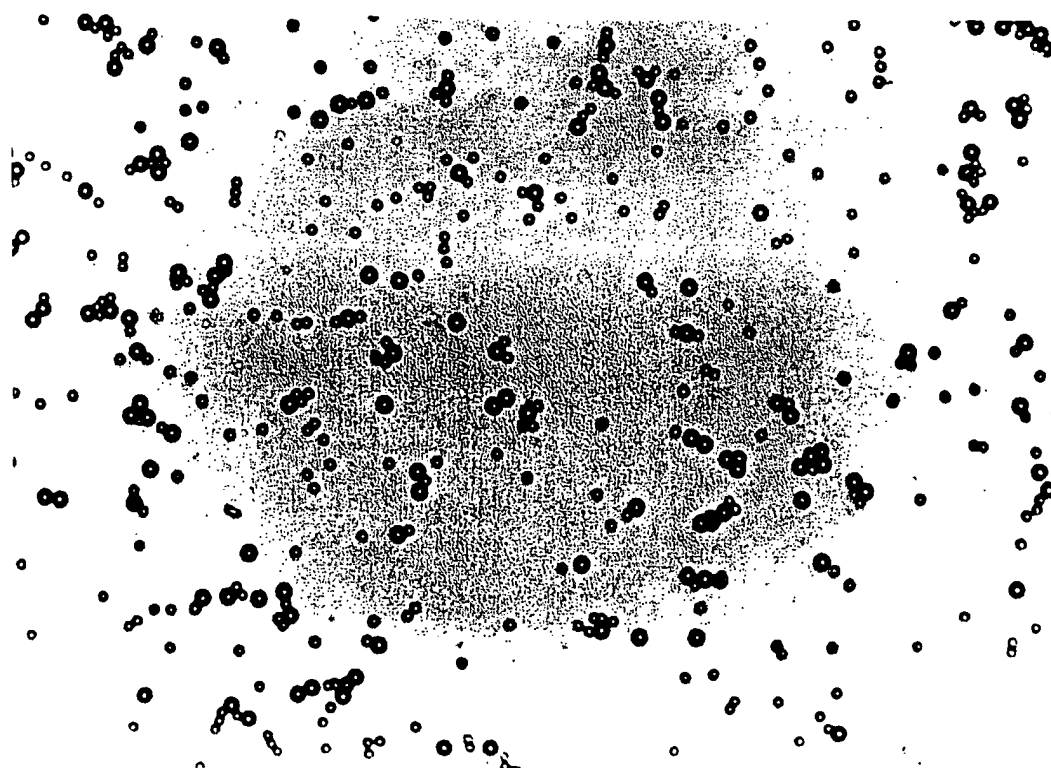


FIG. 6B



4/6

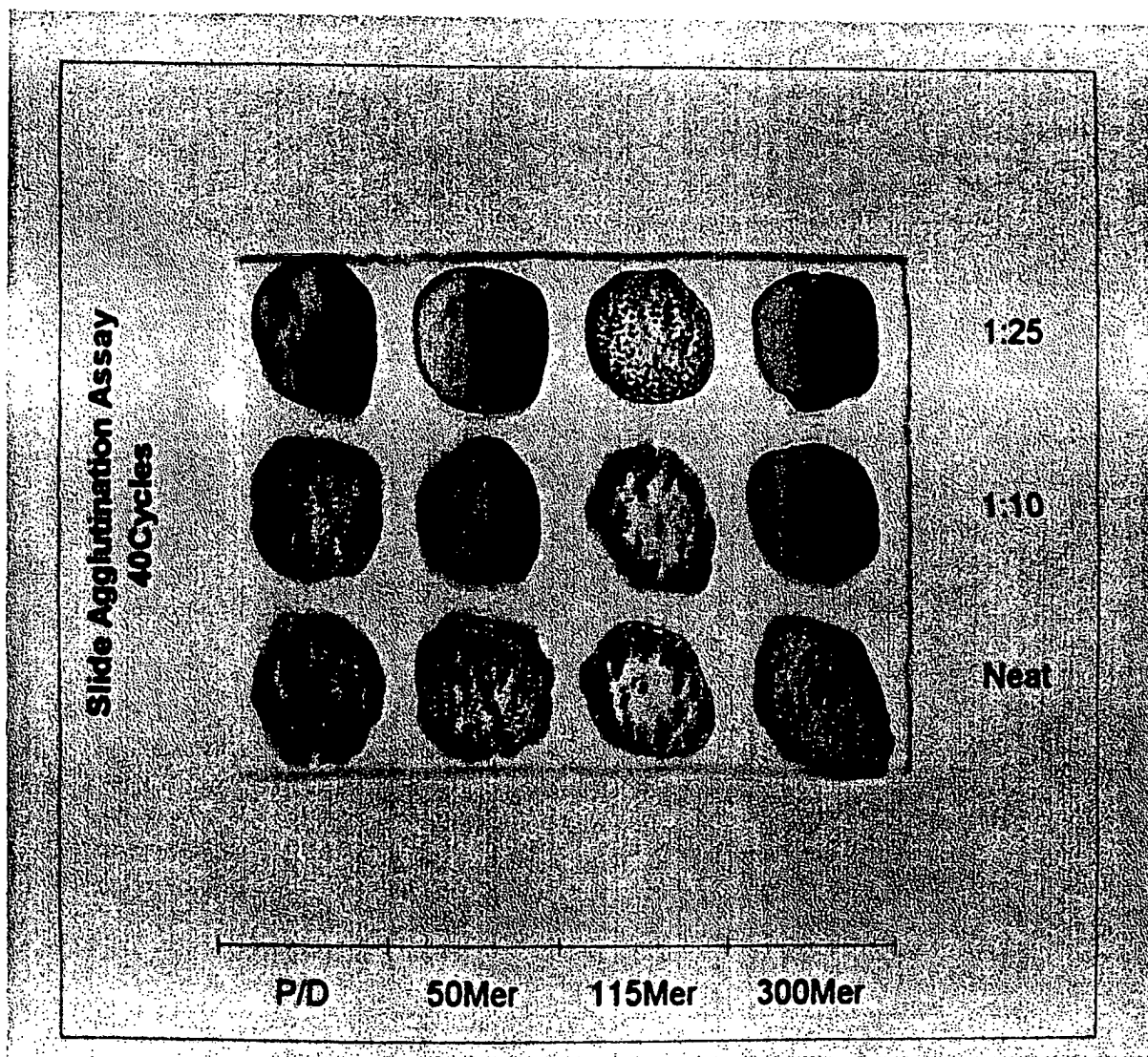


FIG. 7

5/6

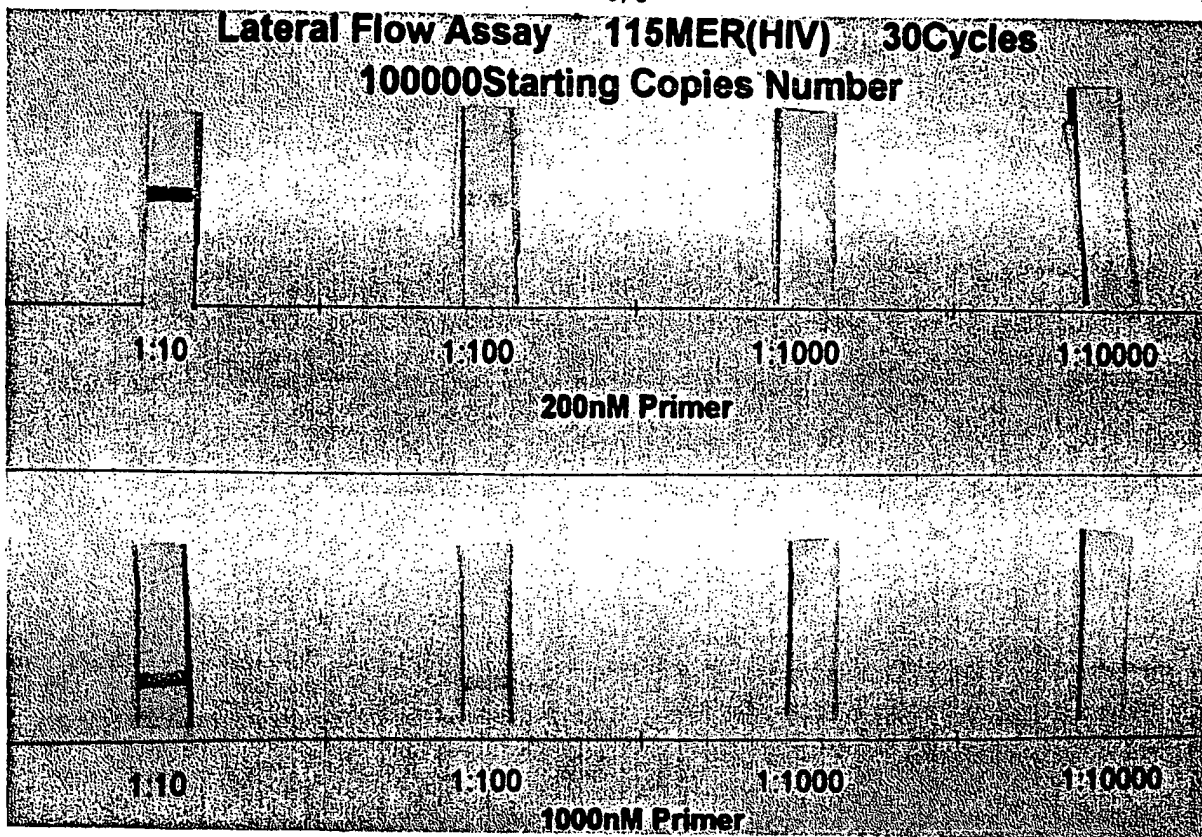


FIG. 9A

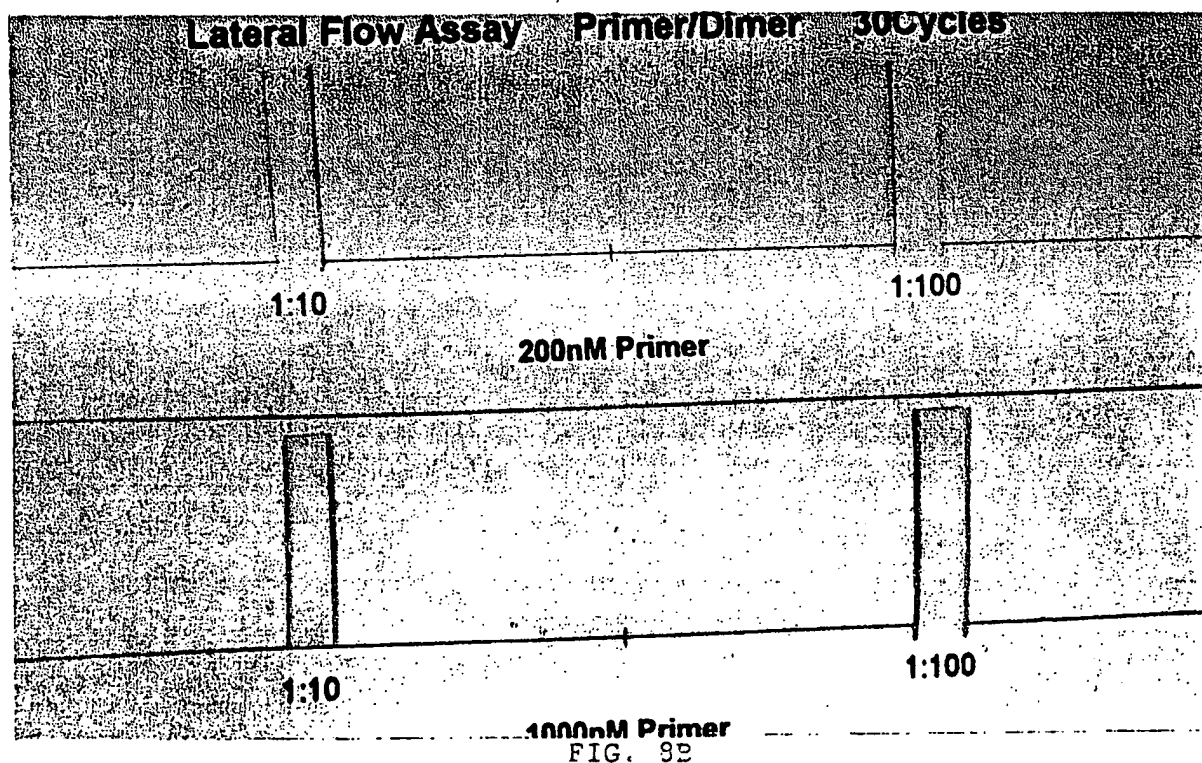


FIG. 9B

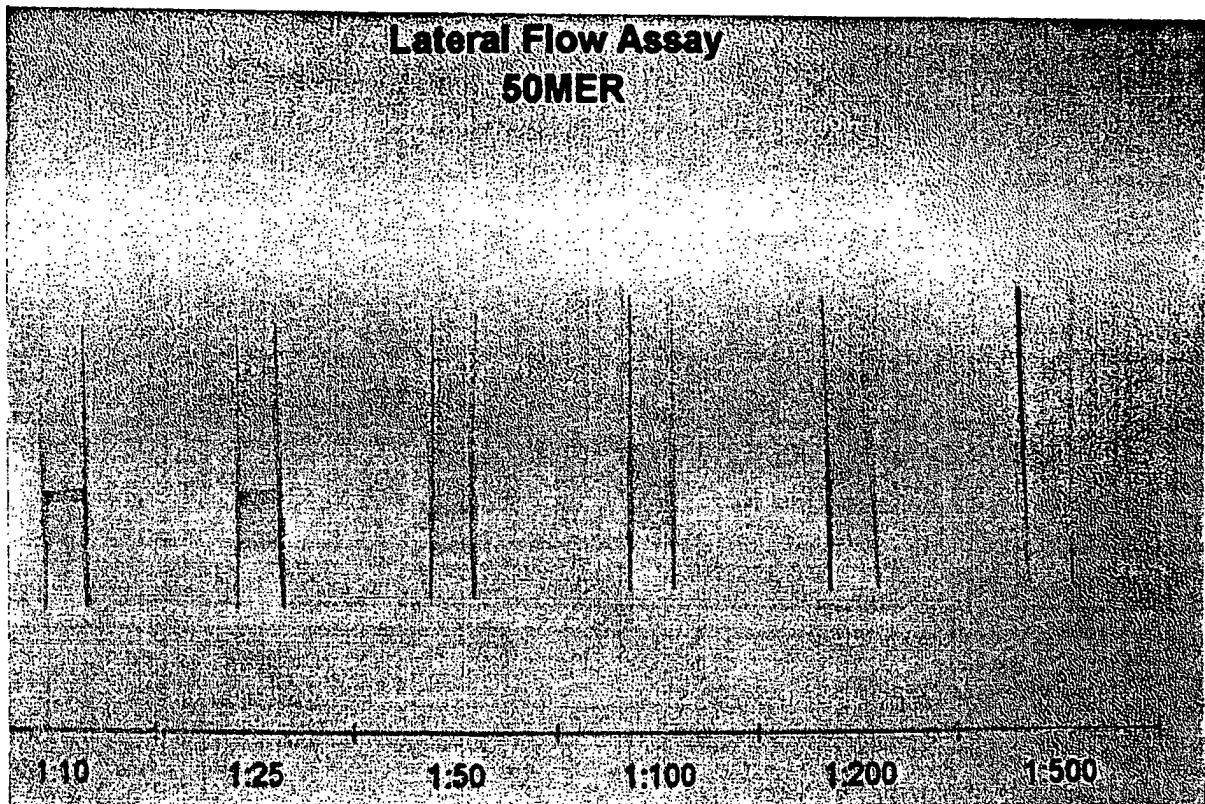


FIG. 9

# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US96/11619

## A. CLASSIFICATION OF SUBJECT MATTER

IPC(6) :C12P 19/34; C12Q 1/68, 1/70; G01N 33/53; C07H 21/04; C07K 16/00

US CL :435/ 91.2, 6, 5, 7.1, 7.9; 536/ 24.3, 24.32, 24.33; 530/388.1

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 435/ 91.2, 6, 5, 7.1, 7.9; 536/ 24.3, 24.32, 24.33; 530/388.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

Please See Extra Sheet.

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US, A, 5,415,839 (ZAUN ET AL.) 16 May 1995, column 1, lines 63-70, column 32, lines 21-70, column 34, lines 20-35, column 35, lines 61-70, column 36 lines 1-10, column 37 lines 5-26.	1-31
Y, P	US, A, 5,489,507 (CHEHAB) 06 February 1996, column 4, lines 5-33 , column 5, lines 22-46.	1-31
Y, E	US, A, 5,543,305 (CUMMINS ET AL.) 06 August 1996, column 4, lines 41-53, column 7, lines 24-35, column 8, lines 59-70.	6,8-10,12, 26- 31
Y, E	US, A, 5,541,069 (MORTENSON ET AL.) 30 July 1996, column 1, column 8, line 57.	19, 20, 26-31



Further documents are listed in the continuation of Box C.



See patent family annex.

* Special categories of cited documents:	*T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
*A* document defining the general state of the art which is not considered to be of particular relevance	*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
*E* earlier document published on or after the international filing date	*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
*L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*G* document member of the same patent family
*O* document referring to an oral disclosure, use, exhibition or other means	
*P* document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search

03 SEPTEMBER 1996

Date of mailing of the international search report

02 OCT 1996

Name and mailing address of the ISA/US  
Commissioner of Patents and Trademarks  
Box PCT  
Washington, D.C. 20231

Facsimile No. (703) 305-3230

Authorized officer

DIANNE REES

Telephone No. (703) 308-0196

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US96/11619

## B. FIELDS SEARCHED

Electronic data bases consulted (Name of data base and where practicable terms used):

APS, BIOSIS, BIOTECHABS, BIOTECHDS, BIOBUSINESS, CABA, CAPLUS, CANCERLIT, DISSABS, EMBASE, IFIPAT, JAPIO, LIFESCI, MEDLINE, USPATFULL, WPIDS.

search terms: primers, PCR, amplification, agglutination, haptens, receptors, binding pairs, ligands, biotin, digoxigenin. FAM, microparticles, steroids, halogens, 2,4-nitrophenyl, Cytomegalovirus, HIV, Mycobacterium tuberculosis, isothermal amplification, lateral flow membrane assay, filtration, microporous, nitrocellulose, nylon, latex, polyethylene, polypropylene, polymethacrylate, polystyrene.